IUBMB Enzyme Nomenclature

EC 1.3.1.2

Common name: dihydropyrimidine dehydrogenase (NADP)

Reaction: 5,6-dihydrouracil + NADP = uracil + NADPH,

Other name(s): dihydrothymine dehydrogenase; dihydrouracil dehydrogenase (NADP); 4,5-dihydrothymine: oxidoreductase; DPD; DHPDH; dehydrogenase, dihydrouracil (nicotinamide adenine dinucleotide phosphate); dihydrouracil dehydrogenase (NADP); DHU dehydrogenase; hydropyrimidine dehydrogenase

Systematic name: 5,6-dihydrouracil:NADP 5-oxidoreductase

Comments: Also acts on dihydrothymine.

Links to other databases: BRENDA, EXPASY, KEGG, WIT, CAS registry number: 9029-01-0

References:

- 1. Fritzson, P. Properties and assay of dihydrouracil dehydrogenase of rat liver. *J. Biol. Chem.* 235 (1960) 719-725.
- 2. Shiotani, T. and Weber, G. Purification and properties of dihydrothymine dehydrogenase from rat liver. *J. Biol. Chem.* 256 (1981) 219-224. [Medline UI: 81093928]

[EC 1.3.1.2 created 1961, modified 1986]

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(54) Title: DIHYDROPYRIMIDINE DEHYDROGENASE COMPOSITIONS AND METHODS OF TISE	(74) Agent: WILSON, Mark, B.; Arnold, White & Durkee, P.O. Box 4433, Houston, TX 77210-4433 (US).	(72) Inventors: DIASIO, Robert, B.; 1225 Branchwater Lane, Birmingham, AL 35216 (US), LU, Zhihong; 1824 Russet Woods Lane, Birmingham, AL 35244 (US). ZHANG, Ruiwer, 1824 Russet Woods Lane, Birmingham, AL 35244 (US). JOHNSON, Martin; 211 Coral Circle, Alabaster, AL 35007 (US). CHENG, Xiaogang; 840 - 16th Street South, Birmingham, AL 35205 (US).	(71) Applicant: THE UAB RESEARCH FOUNDATION (US/US); Suite 1120G, 701 South 20th Street, Birmingham, AL 35294-0111 (US).	(30) Priority Data: 08/227,357 13 April 1994 (13.04.94)	(22) International Filing Date: 13 April 1995 (13.04.95)	(21) International Application Number: PCT/US95	C12N 15/53, 9/02, C12Q 1/32, C07K 16/40	(51) International Patent Classification 6:	INTERNATIONAL APPLICATION PUBLISH
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MIONE AND METHODS OF IRE			Published With international search report.	TI, TT, UA, UZ, VN, European patent (AT, BR, CH, DE, DK, BS, FR, GB, GR, IE, IT, LU, MC, UI, FT, SS), OAP! patent (BF, BJ, CF, CG, CI, CM, GA, GN, ML, MR, NE, SN, TD, TG), ARIPO patent (KE, MW, SD, SZ, UG).	11 1995 (13.04.95) KG, KP, KR, KZ, LK, LR, LT, LU, LV, MD, MG, MN, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK,	(81) Designated States: AM, AT, AU, BB, BG, BR, BY, CA, CH,	(43) International Publication Date: 26 October 1995 (26.10.95)	(11) International Publication Number: WO 95/28489	INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCI)

(54) Time: DIHYDROPYRIMIDINE DEHYDROGENASE COMPOSITIONS AND METHODS OF USE

(57) Abstract

Disclosed are methods and compositions for use in detecting and quantifying the enzyme dihydropyrimidine dehydrogenase (DPD) for use in, e.g., optimizing 5-fluorouracil doces given to cancer patients. Particularly described are antibodies, including monoclonal antibodies, to the human form of DPD; DNA sequences from bovine and human DPD; immunological and molecular biological means by which to detect DPD; and methods of designing effective cancer treatment strategies based upon information gained concerning DPD levels. Also disclosed is molecular characterization of a genetic leation leading to DPD deficiency in humans and diagnostic methods for genetic screening of this mutation for patients undergoing FUra treatment.

2. DNA Sequence Analysis of DPD Gene in a DPD-Deficient Patient

position 2894. Translation of the cDNA demonstrated that this resulted in a additional single nucleotide difference from that of control; A (control) to T (deficient) at Complete sequence analysis of the DPD deficient patient's cDNA also revealed an

5 normal DPD activity but was identified exclusively in the DPD deficient patient's cDNA. adenosine deletion resulting in a frameshift was not found in any individuals having common in the general population and may represent an allelic variant. In contrast, the individuals having normal DPD activity demonstrated that this nucleotide substitution was sequence analysis of multiple PCR[™] reactions flanking this region from a number of nonconservative amino acid substitution (Asp to Val). Subsequent subcloning and

20 ᇙ other identical to normal), present in approximately equal amounts. The identification of analysis of several clones (from multiple PCR" reactions) from the deficient patient containing the sequence of interest (FIG. 7A, FIG. 7B, FIG. 7C, and FIG. 7D). Sequence this patient is heterozygous for the single base deletion. both the normal and mutant allele (adenosine deletion) in the genomic DNA confirm that indicated the presence of two different alleles (one of these containing the deletion, the confirm that this patient was heterozygous for this mutation. Primers were designed based on the cDNA sequence to amplify a 573 base pair DNA fragment from the exon patient (two out of the ten subclones), studies were undertaken with genomic DNA to Since this deletion was initially identified in the cDNA of the DPD deficient

25 explanation for reduced DPD activity. This frameshift has also been identified in an represents the first molecular characterization of a DPD deficient patient, and provides an genomic DNA has demonstrated that this patient is heterozygous for this mutation. This translation at codon 335 generating a 36,500 dalton protein. Analysis of the patient's patient contains an adenosine deletion that causes a frameshift resulting in truncation of additional unrelated DPD deficient patient who also exhibited severe FUra toxicity. In summary, the gene and the poly(A)+ RNA encoding the OPD protein in this